

18. Hamilton, C.M. (1997). A binary-BAC system for plant transformation with high-molecular-weight DNA. *Gene* 200:107-116.
19. Hengen, P. N., and Iyer, V. N. (1992). DNA cassettes containing the origin of transfer (*oriT*) of two broad-host-range transfer systems. *BioTech.* 13:56-62.
20. Hoekema, A. et al. (1983). A binary plant vector strategy based on separation of *vir*- and T-region of the *Agrobacterium tumefaciens* Ti-plasmid. *Nature* 303:179-180.
21. Ingersoll, J.C., Rothenberg, M., Liedl, B.E., Folkerts, K., Garvin, D., Hanson, M.R., Doyle, J.J., and Mutschler, M.A. (1994). A novel anther-expressed *adh*-homologous gene in *Lycopersicon esculentum*. *Plant Molec. Biol.* 26:1875-1891.
22. Jones, J. D. G. et al. (1992). Effective vectors for transformation, expression of heterologous genes, and assaying transposon excision in transgenic plants. *Transgenic Research* 1:285-297.
23. Jouanin, L. et al. (1985). Localization and restriction maps of the replication origin regions of the plasmids of *Agrobacterium rhizogenes* strain A4. *Mol. Gen. Genet.* 201:370-374.
24. Keasling, J.D. (1999). Gene-expression tools for the metabolic engineering of bacteria. *Tibtech* 17:452-460.
25. Knauf, V. C., and Nester, E. W. (1982). Wide host range cloning vectors: a cosmid clone bank of an *Agrobacterium* Ti plasmid. *Plasmid* 8:45-54.
26. Kohara, Y. et al. (1987). The physical map of the whole *E. coli* chromosome: application of a new strategy for rapid analysis and sorting of a large genomic library. *Cell* 50:495-508.

27. Kroll et al. (1998). Natural genetic exchange between *Haemophilus* and *Neisseria*: intergeneric transfer of chromosomal genes between major human pathogens. *Proc. Nat'l. Acad. Sci. (USA)* 95:12381-12385.
28. Lawrence, J. (1999). Selfish operons: the evolutionary impact of gene clustering in prokaryotes and eukaryotes. *Curr. Opinion in Genetics and Development* 9:642-648.
29. Leong, S. A. et al. (1982). Heme biosynthesis in *Rhizobium*. Identification of a cloned gene coding for S-aminolevulinic acid synthetase from *Rhizobium meliloti*. *J. Biol. Chem.* 257:8724-8730.
30. Lewin, A, D. Jacob, B. Freytag, B. Appel. (1998). Gene expression in bacteria directed by plant-specific regulatory sequences. *Transgenic research* 7:403-411.
31. Lin-Chao, S. et al. (1992). High copy number of the pUC plasmid results from a Rom/Rop-suppressible point mutation in RNAII. *Molec. Microbiol.* 6:3385-3393.
32. Lopilato, J. et al. (1986). Mutations in a new chromosomal gene of *Escherichia coli* K-12, *pcnB*, reduce plasmid copy number of pBR322 and its derivatives. *Mol. Gen. Genet.* 205:285-290.
33. Low, K. B. (1972). *Escherichia coli* K-12 F-prime factors old and new. *Bacteriol. Rev.* 36:587-607.
34. Miranda, A. et al. (1992). *Agrobacterium tumefaciens* transfers extremely long T-DNAs by a unidirectional mechanism. *J. Bact.* 174:2288-2297.

35. Mozo, T., and Hooykaas, P. J. J. (1991). Electroporation of megaplasmid into *Agrobacterium*. Plant Mol. Biol. 16:917-918.
36. O'Conner, M. et al. (1989). Construction of large DNA segments in *Escherichia coli*. Science 244:1307-1312.
37. Osoegawa et al. (1998). An improved approach for construction of bacterial artificial chromosome libraries, Genomics 52:1-8.
38. Pace, N.R. (1996). A molecular view of microbial diversity and the biosphere. Science 276:734-740.
39. Peralta, E. G., and Ream, L. W. (1985). T-DNA border sequence required for crown gall tumorigenesis. Proc. Natl. Acad. Sci. USA. 82:5112-5116.
40. Peralta, E. G. et al. (1986). Overdrive, a T-DNA transmission enhancer on the *A. tumefaciens* tumor-inducing plasmid. EMBO J. 5:1137-1142.
41. Pierce, J. C. et al. (1992). A positive selection vector for cloning high molecular weight DNA by the bacteriophage P1 system: Improved cloning efficacy. Proc. Natl. Acad. Sci. USA. 89:2056-2060.
42. Rackwitz, H. R. et al. (1985). Analysis of cosmids using linearization by phage lambda terminase. Gene 40:259-266.
43. Richard-L. Qin-L-X. Goldberg-R., (1996). Clustered genes within the genome of *Arabidopsis thaliana* encoding pectin methylesterase-like enzymes. Gene 170:207-211.